

REMARKS

Claims 5-11 and 18-35 are pending in this application. Claims 1-4 and 12-17 have been cancelled without prejudice or disclaimer. Claims 18-35 have been added. Claims 5-8 have been amended. Claim 5 was amended to depend from new claim 18 and to delete reference to SEQ ID NO:3. Claim 6 was amended to recite "detecting hybridization between a nucleic acid comprising at least 10 contiguous nucleotides of SEQ ID NO:1 and a PCGEM1 nucleic acid." Support for this amendment can be found throughout the specification including, for example, at page 14. Claims 7 and 8 were amended to more accurately reflect their dependency from claims 6 and 7, respectively.

Support for new claims 18-35 can be found throughout the specification, including, for example, at pages 13-16. Thus, no new matter is added by this amendment.

The application has also been amended to insert the required reference to the parent application(s), for which benefit is claimed. It has also been amended to insert the Sequence Listing before the claims. Applicants have filed concurrently with this continuation application a Request for Transfer of Computer Readable Form Under 37 C.F.R. § 1.821(e).

Applicants also request that Figure 8 be amended as indicated in the attached Annotated Sheet (Attachment B). The same change was made to Figure 8 in the parent application. Specifically, nucleotide 302 of SEQ ID NO:1 has been changed from a thymine base ("T") to a guanine base ("G"). As explained during prosecution of the

Rule 1.53(b) continuation appln. of
Serial No. 09/534,072

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parent application, this discrepancy at nucleotide 302 arose from a sequencing error,
and the amendment correcting this discrepancy is fully supported by the specification.

Please grant any extensions of time required to enter this paper and charge any
additional required fees to our deposit account 06-0916.

Respectfully submitted,

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Dated: March 18, 2004

By: 

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